

Figure 1

trypsin: domain 1 of 1, from 217 to 443: score 293.0, E = 3.2e-92

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*->IvGGreaqpgsfsgPwqvslqvrsgggsrkhfCGGSLisenwVLTAA
IvGG + ++ Pwq+slq+ + +h+CGGs+i + w++TAA
14094 217 IVGGNMSLLSQW--PWQASLQFQG-----YHLCGGSVITPLWIITAA 256

HCvsgaasapassvrVSlsvrlGehnlsltegteqkfdvkkttiivHpnyn
HCv++ +++s+++ +G +sl + v+k i+ H++y
14094 257 HCVYD--LYLPKSWTI----QVGL--VSLLDNPAPSHLVEK-IVYHSKYK 297

pdtldngaYdnDiAlLkLkspgvtlgtvvpicLpsassdlpvGttctvs
p+ l+ nDiAl+kL++p +t+++ ++p+cLp + ++p+G c+ s
14094 298 PKRLG-----NDIALMKLAGP-LTFNEMIQPVCLPNSEENFPDGKVCWTS 341

GwGrrptknlg.lsdLqevvvpvsretCrsayeyggtdDkvefvtdnm
GWG t+++g+ s +L ++ vp++s++ C+++ ygg +++ m
14094 342 GWGA--TEDGGdASPVLNHAAPVLISNKICNHRDVGIGI-----ISPSM 383

iCagal.ggkdaCqGDSGGPLvcsgdnrdgrwelvGivSwGsygCargnk
+Cag+l+gg+d+CqGDSGGPLvc w+lvG +S+G gCa+ nk
14094 384 LCAGYLtGGVDSCQGDSSGGLVQCER---RLWKLVGATSFSG-IGCAEVNK 429

PGvytrVssyldWI<-*
PGvytrV+s+ldWI
14094 430 PGVYTRVTSFLDWI 443

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Fig. 2A

trypsin_2: domain 1 of 1, from 216 to 443: score 328.2, E = 9.2e-95

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*->RIVGGseakigsfPwqvslq.....CGGSLisprwVLTAAHC.....
RIVGG+ + ++Pwq+slq ++ + CGGS+I+p w++TAAHC +
14094 216 RIVGGNMSLLSQWPWQASLQfagyhLCGGSVITPLWIITAAHCvydl 262

.....rVrlGshdlssgeeteggprldspggqvikVskiiievHpnYn..
+++ ++ +G +l + + + V+ki+ H +Y ++
14094 263 ylpkswTIQVGLVSLDNP-----APSHLVEKIV-YHSKYKpk 299

...NDIALlLkLkepvtlsdsntvrPicLPssneiktsegntvpaGttctv
+ +NDIAL+kL+ p+t+++ ++P+cLP+s +++p+G c+
14094 300 rlgNDIALMKLAGPLTFNE--MIQPVCLPNS-----EENFPDGKVCWT 340

sGWGrtsegpeesgggslpdvLqevnvpivsnetCr.....Ml
sGWG t++ gg + vL ++ vp++sn+ C++++ +++ ++Ml
14094 341 SGWGATED-----GGDASPVLNHAAPVLISNKICNhrdvgyggiispsML 384

CAGyleggntpgGkDaCqGDSGGPLvc.....vLvGiVSWGssslgCa
CAGyl+ gG+D+CqGDSGGPLvc++++ ++LVG +S+G +gCa
14094 385 CAGYLT-----GGVDSCQGDSSGGLVQCqerrlWKLVGATSFSG----IGCA 425

rpnkPGVYTrVssyldWI<-*
+ nkPGVYTrV+s+ldWI
14094 426 EVNKPVGYYTRVTSFLDWI 443

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Fig. 2B

Fig. 3A

Begin SEQ ID NO 6

End SEQ ID NO 6

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*->stCggpdeFqCgsgrrCIprswvCDGdpDCeDGSDEslenCaa<-*
+C+ ++++C+s+ CI +CDG DC+DG+DE +C++
14094 71 FDCS--GKYRCRSSFKCIELIARCDGVSDCKDGEDE--YRCVR 109
  
```

Fig. 3B

Begin SEQ ID NO 7

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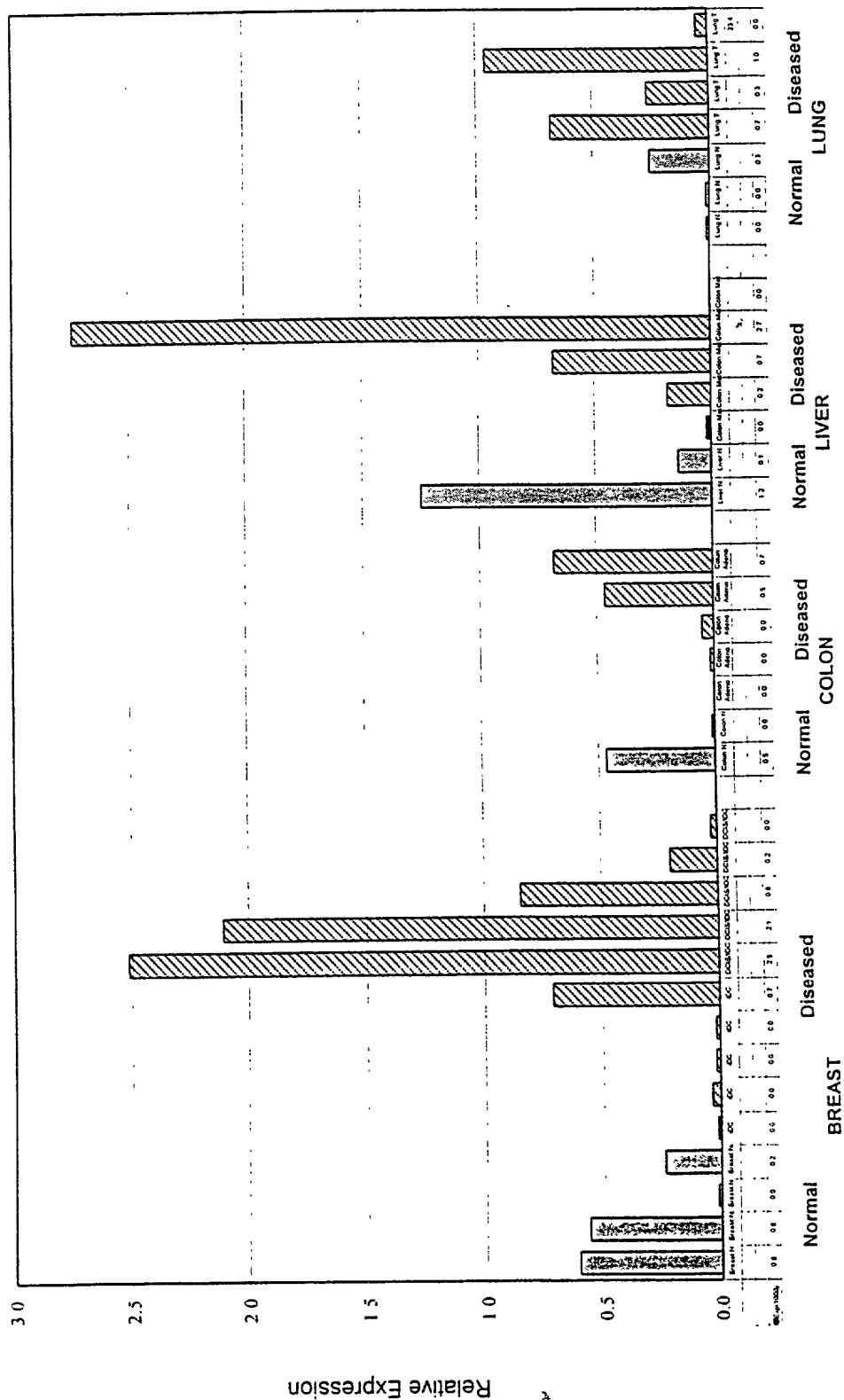
*->vgGssrCeGrVEVrhgskwgtVCdssWslrdanvdpQaskvCrqLG
vgG +++ +++V+ + W+t C+++W + anv +C+qLG
14094 110 VGG--QNA-VLQVF-TAASWKTMCSDDWKGHYANV-----ACAQLG 146

CGgavsl1.gpyfsegggPagqreiwldgvnCsGnE...tsLsqCpvrvt
+ vs+ + s+ g ++++++ + +++++ +
14094 147 FPSYVSSDnLRVSSLEG-----QFREEFVSIDHLlpdDKVTALHHS-- 187

ppglrqqcshdgedagVvCs<-*
++ ++c g+ ++C
14094 188 -VYVREGCAS-GHVVTLQCT 205
  
```

END SEQ ID NO 7

Figure 4



14094 Expression in Clinical Ovary Samples

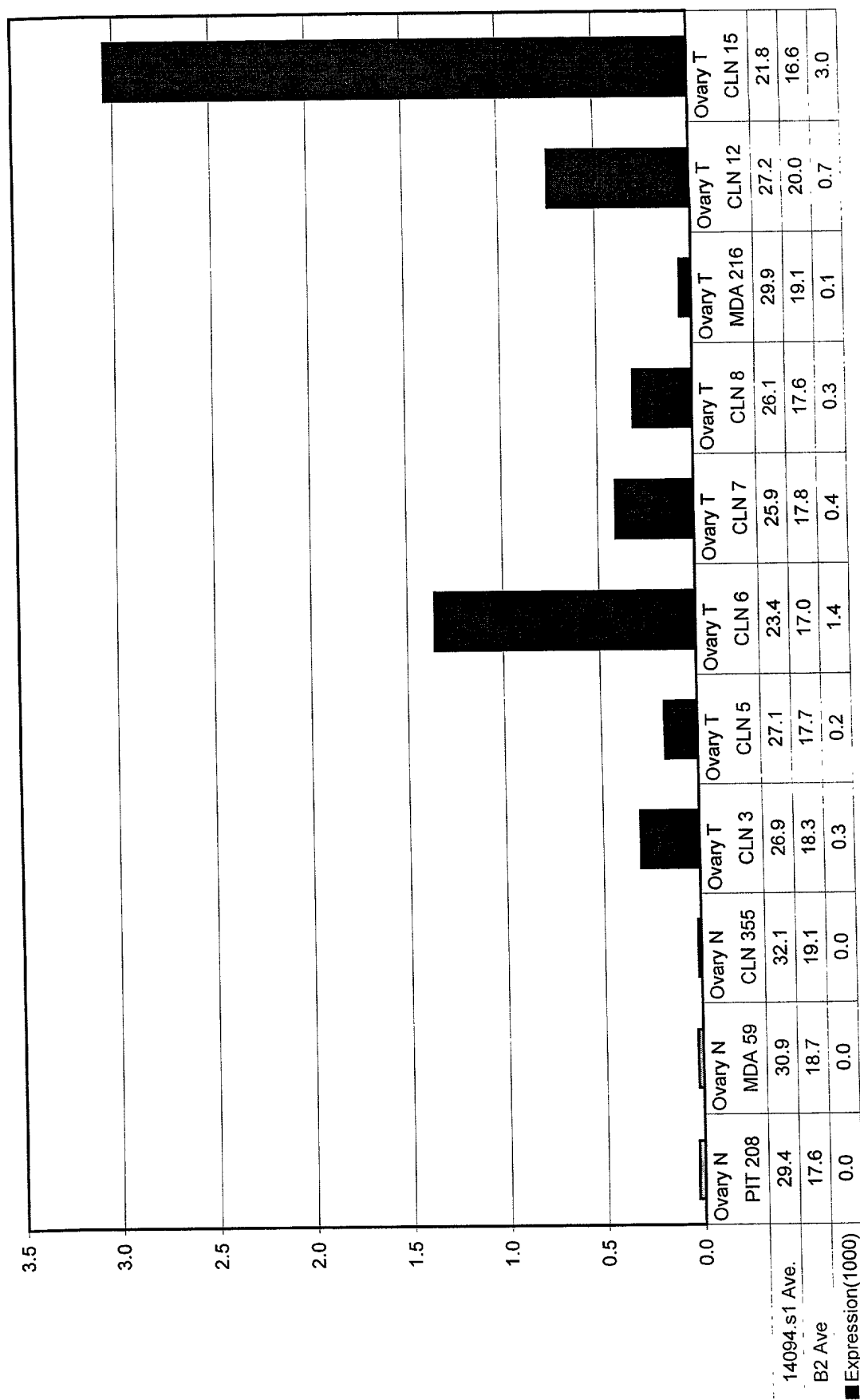


Figure 5

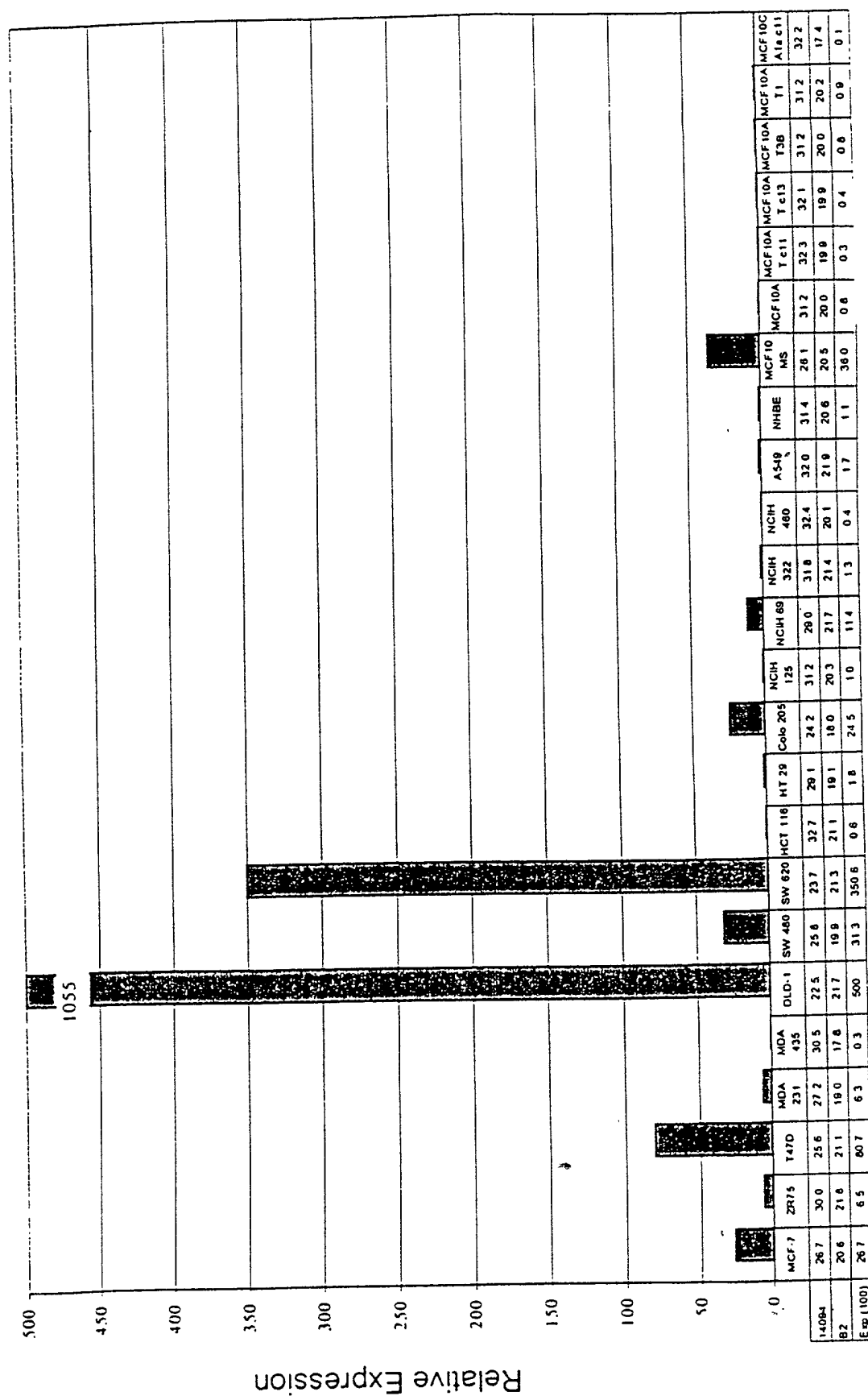


Fig. 6